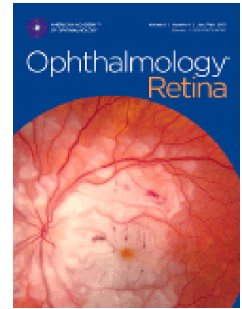


# Journal Pre-proof



*CERKL*-associated retinal dystrophy: Genetics, Phenotype and Natural History

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1 CERKL-associated retinal dystrophy: Genetics, Phenotype and Natural History

2

3 **Running head:** CERKL-associated retinal dystrophy

4

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42

**Abstract**

**Purpose:** To analyze the clinical characteristics, natural history, and genetics of *CERKL*-associated retinal dystrophy in the largest series to date.

**Design:** Multicenter retrospective cohort study.

**Subjects:** 47 patients (37 families) with likely disease-causing *CERKL* variants

**Methods:** Review of clinical notes, ophthalmic images, and molecular diagnosis from two international centres.

**Main outcome measures:** Visual function, retinal imaging and characteristics were evaluated and correlated.

**Results:** The mean age at the first visit was  $29.6 \pm 13.9$  years and the mean follow-up time was  $9.1 \pm 7.4$  years. The most frequent initial symptom was central vision loss (40%) and the most common retinal feature was well-demarcated areas of macular atrophy (57%). Seventy percent of the participants had double-null genotypes and 64% had electrophysiological assessment. Amongst the latter, 53% showed similar severity of rod and cone dysfunction, 27% revealed a rod-cone, 10% a cone-rod, and 10% a macular dystrophy dysfunction pattern. Patients without double-null genotypes tended to have fewer pigment deposits and included a higher proportion of older patients with a relatively mild electrophysiological phenotype.

Longitudinal analysis showed that over half of the cohort lost 15 ETDRS letters or more in at least one eye during the first 5 years of follow up.

**Conclusions:** The phenotype of *CERKL*-retinal dystrophy is broad, encompassing isolated macular disease to severe retina-wide involvement, with a range of functional phenotypes, generally not fitting in the rod-cone/cone-rod dichotomy. Disease onset is often earlier, with more severe retinal degenerative changes and photoreceptor dysfunction, in nullizygous cases.



## 66 Introduction

67 Inherited retinal diseases (IRDs) are a group of heterogeneous stationary or progressive retinal  
68 disorders, that often cause visual impairment.<sup>1</sup> IRDs represent a leading cause of visual disability  
69 in the working age population in the UK.<sup>2</sup> Depending on whether rod or cone photoreceptors are  
70 primarily (or more severely) affected, two main phenotypes of IRD can be differentiated – cone-  
71 rod (CORD) and rod-cone dystrophy (RCD); the latter also known as retinitis pigmentosa (RP).  
72 Macular dystrophy (MD) corresponds to an IRD where dysfunction is confined to the macula.  
73 Halting the progression of these conditions and alleviating their symptoms is the subject of  
74 multiple avenues of research.

75 Over 300 genes and loci are currently known to cause IRD (<https://web.sph.uth.edu/RetNet/>).  
76 Ceramide-kinase like (*CERKL*, MIM 608381) is amongst the rare causative genes in the UK,  
77 detected only in 17 families of a large cohort of over 3000 genetically diagnosed IRD families by  
78 2020.<sup>3</sup> It is also reported to be an uncommon cause of CORD in a German cohort.<sup>4</sup> Other groups  
79 however, report a higher prevalence of *CERKL*-associated retinopathy, accounting for 33% and  
80 5% of autosomal recessive IRD in Yemenite Jewish and Spanish populations, respectively.<sup>5,6</sup>  
81 *CERKL* was associated with 8% of IRD cases in Tunisia,<sup>7</sup> and was one of the commonest genes  
82 in a large Spanish RP cohort study.<sup>8</sup> To date, four founder disease-causing variants have been  
83 described; NM\_001030311.2: c.847C>T (p.Arg283Ter) (reported as NM\_201548.5: c.769C>T)  
84 from Spain, c.238+1G>A from the Yemenite Jewish population, c.375C>G (p.Cys125Trp) from  
85 the Finnish, and c.365T>G (p.Leu122Arg) from indigenous African populations in South  
86 Africa.<sup>5,9-11</sup>

87 *CERKL* was first associated with IRD in 2004, after Tuson *et al.* analyzed RP26, a 17.4 Mb locus  
88 in chromosome 2 known to cause RP in Spanish families.<sup>9</sup> It is a 13-exon gene that encodes a  
89 532-amino acid protein, with a diacylglycerol kinase and a pleckstrin homology domains.<sup>12</sup> It is  
90 widely expressed in the kidney, lung, skin, and pancreas; and it is known to have a protective  
91 role against oxidative stress-induced apoptosis.<sup>13</sup> It also interferes with mitochondrial  
92 metabolism, stress granules, and autophagy regulation; yet its pathophysiology remains to be  
93 fully elucidated.<sup>14</sup>

94 Biallelic loss-of-function variants in *CERKL* have been associated with both RCD and  
95 CORD.<sup>15,16</sup> However, the reported retinal phenotype in both cases appears to be somewhat

96 similar: a widespread retinopathy with early maculopathy and a similar severity of rod and cone  
97 dysfunction electrophysiologically.<sup>5,15</sup> Patients often report concurrent cone- (e.g., central vision  
98 and colour discrimination impairment) and rod-related (e.g., nyctalopia, peripheral field loss)  
99 symptoms, becoming noticeable from adolescence/young adulthood, with an unclear timeline.<sup>17</sup>  
100 *CERKL* can also present with a Stargardt-like phenotype, often being a differential diagnosis  
101 when no variants are found in *ABCA4*. Nevertheless, there remains a paucity of detailed  
102 phenotypic characterization in a large cohort, thereby limiting our understanding of *CERKL*-  
103 associated retinopathy and its natural history. This study aims to establish the phenotypic and  
104 genetic spectrum by examining the largest case series of *CERKL*-associated retinopathy patients  
105 to date, to better understand this disorder and to optimize future clinical management.

106

## 107 **Methods**

108 This study was a retrospective, consecutive case series of patients who attended Moorfields Eye  
109 hospital (MEH, London, UK) and the Royal Victoria Eye and Ear Hospital (Dublin, Ireland)  
110 with a retinal dystrophy, and were found to have likely disease-causing variants in *CERKL*. At  
111 MEH, patients were identified through the inherited eye disease database. Informed consent was  
112 obtained from all patients. Ethical approval was provided by the local ethics committee and the  
113 study honored the tenets of the Declaration of Helsinki.

114 Relevant patient data was retrieved from electronic healthcare records, case notes, and imaging  
115 software systems. Snellen visual acuities were recorded and converted to LogMAR for the  
116 purpose of statistical analysis. Count fingers vision was given a value of LogMAR 1.98, hand  
117 motion LogMAR 2.28, light perception LogMAR 2.7, and no light perception LogMAR 3.<sup>18</sup>  
118 Asymmetric best-corrected visual acuity (BCVA) was defined as a difference  $\geq 0.3$  LogMAR  
119 (equivalent to 15 Early Treatment Diabetic Retinopathy Study -ETDRS- letters) between eyes.  
120 Patients were categorized using the World Health Organization (WHO) visual impairment  
121 criteria, that defines no or mild visual impairment as  $BCVA \leq 0.48$  (6/18, 20/60), moderate  
122 impairment as  $BCVA > 0.48$  and  $\leq 1.0$  (6/60, 20/200), severe as  $BCVA > 1.0$  and  $\leq 1.3$  (3/60,  
123 20/400), and blindness as  $BCVA > 1.3$ . Records of visual field were limited within our cohort;  
124 therefore, we classified patients according to BCVA only. 'Low vision' corresponds to patients  
125 with moderate and severe visual impairment.

126 Further clinical assessment consisted of dilated fundus examination, spectral-domain optical  
127 coherence tomography (SD-OCT, Heidelberg Spectralis, Heidelberg Engineering, Inc.,  
128 Heidelberg, Germany), fundus autofluorescence (Heidelberg Spectralis, Heidelberg Engineering,  
129 Inc., Heidelberg, Germany and Optos PLC, Dunfermline, UK) and ultrawide field fundus  
130 pseudocolor photography (Optos PLC). OCT thickness in the general population was extracted  
131 from Invernizzi *et al.*<sup>19</sup> Fovea-centered macular volume scans were performed in a 6-mm<sup>2</sup> area  
132 that included the 1-, 3-, and 6-mm grid templates from the ETDRS. Inner limiting membrane and  
133 Bruch membrane were automatically segmented by the manufacturer software (Heyex version  
134 1.9.14.0; Heidelberg Engineering) and adjusted manually as needed by a trained ophthalmologist  
135 (M.D.V.). Ellipsoid zone (EZ) width and outer nuclear layer thickness (ONLT) were measured  
136 manually at the foveal scan. Patients with macular cysts, oedema, and/or only line scans were  
137 excluded from quantitative assessment. Non-ocular issues were defined as those commonly  
138 found in syndromic IRDs, such as musculoskeletal, renal, cardiac, audiological, or  
139 neuro/psychological.<sup>20</sup>

140 Pattern and full-field electroretinogram (PERG; ERG) testing was performed incorporating the  
141 International Society for Clinical Electrophysiology of Vision (ISCEV) standards.<sup>21</sup> Pattern ERG  
142 P50 was used as an objective measure of macular function and full-field ERG was used to assess  
143 generalized (mainly peripheral) rod and cone system function. The main dark-adapted (DA) and  
144 light-adapted (LA) ERG components were quantified and compared with age-matched control  
145 data from healthy subjects (age range: 10-79 years).<sup>22</sup> ERG amplitudes were plotted as a  
146 percentage of the age-matched lower limit of the reference range or as a difference from the age-  
147 matched upper limit of peak time. The reference limits were defined as lower 5th percentile for  
148 amplitude and 95th percentile for peak times. The full-field ERGs were classified into 4 groups  
149 based on the relative reduction of DA and LA ERGs; rod and cone photoreceptor dysfunction,  
150 rod-cone, cone-rod or predominantly macular dysfunction (normal or near-normal ERGs). For  
151 the purposes of the electrophysiological analysis, a 33% minimum difference in the relative  
152 reduction of DA 10 ERG a-wave and LA 3 ERG b-wave was used to define rod-cone and cone-  
153 rod patterns of dysfunction. An additional patient (age 5 years, ID 41) was tested before his  
154 baseline ophthalmologic assessment according to shorter PERG and ERG protocols using lower  
155 eyelid skin electrodes without mydriasis.

156 DNA was extracted from whole blood and genetic testing was performed using panel-based  
157 targeted next generation sequencing (NGS), whole exome sequencing, or whole genome  
158 sequencing. Where appropriate, blood samples were taken from parents or siblings to confirm  
159 segregation of proposed variants (i.e., to determine if relatives carry one or both variants in one  
160 or two alleles). Other IRD genes were excluded based on coding variant analysis. The  
161 pathogenicity of the variants was determined by implementing the criteria of the American  
162 College of Medical Genetics (ACMG).<sup>23</sup> Likely disease-causing variants correspond to  
163 pathogenic, likely pathogenic, and selected variants of uncertain significance (VUS), based on  
164 family history, phenotype, and/or if concurrent with a pathogenic/likely pathogenic variant. *In*  
165 *silico* molecular genetic analysis was performed for all detected *CERKL* variants (Genome  
166 reference; Hg38 Transcript; NM\_001030311.2, ENST00000339098.9 or NM\_201548.5,  
167 ENST00000410087.8), according to a previous publication.<sup>24</sup> Genotype grouping was performed  
168 according to the presence of null variants (those assumed to result in a loss of function -  
169 nonsense, frameshift, splice site alteration, and exon deletion-); a double-null (DN) genotype  
170 harboring multiple null variants (also known as ‘nullizygous’), and a non-double-null (NDN)  
171 genotype with one or no null variants. Clinical descriptions and parameters were compared  
172 between subgroups of patients with DN and NDN genotypes.

173 GraphPad Prism 8.0.2 (GraphPad Software, San Diego, CA, USA) was used for statistical  
174 analysis. The threshold of significance was set at  $p < 0.05$ . Linear regressions and t-tests were  
175 used for assessment of parametric variables. When testing associations between age and ocular  
176 characteristics, only the right eye was considered to avoid clustering effect. Welch's t-test  
177 variation was employed when the sample sizes were significantly different.

178

## 179 **Results**

### 180 Demographics, phenotype, and visual acuity

181 Forty-seven patients were characterized from 37 pedigrees. Their clinical characteristics are  
182 listed in Table 1 and Supplementary Table 2. Twenty-two individuals were female (47%) and 25  
183 (53%) were male. The mean age  $\pm$  standard deviation (SD) at the first visit was  $29.6 \pm 13.9$  years  
184 (range 8-6 – 67 years) and the median age was 25 years, with 5 participants (11%) having their

185 first visit below the age of 16. The mean follow-up time of the cohort was  $9.1 \pm 7.4$  years and the  
186 mean age at their final visit was  $39.2 \pm 16.6$  years. One patient was deaf and had an autosomal  
187 dominant family history of deafness; no other participant had non-ocular features-issues.

188 Amongst the patients that reported their ethnicity (72%), 19 were of Asian origin, 13 were white  
189 from UK and Europe, and 2 were of African descent. Consanguinity was reported in six families,  
190 where parents were first cousins.

191 Based on ophthalmic history and clinical examination, 26 patients were initially diagnosed with  
192 RP (RCD), 13 with CORD, and 8 with macular dystrophy (MD). The mean age of symptoms  
193 onset was  $19.8 \pm 9$  years (median 17, range 6-47 years), with 16 patients below the age of 16.  
194 Nineteen patients (40%) described central vision loss as their initial symptom, 16 (34%) reported  
195 concurrent blurred central vision and nyctalopia, 5 (11%) reported night blindness first, and the  
196 remaining patients mentioned being initially bothered by scotomas, photophobia, or poor color  
197 discrimination. A prolonged prodromal history of subtle VA decrease was elicited in six patients  
198 (24%), lasting between 1 and 10 years.

199 Mean baseline BCVA was  $0.7 \pm 0.8$  LogMAR OD (median 0.3) and  $0.8 \pm 0.8$  OS (median 0.4).  
200 In those aged 15-24 years (23 participants), BCVA ranged from 0.0 LogMAR to light perception  
201 (LP, mean  $0.5 \pm 0.6$  LogMAR, median 0.3, Figure 1A & B), 18 had none or mild visual  
202 impairment, and 5 had moderate visual impairment. There were 13 patients aged 25-40 years old  
203 whose BCVA ranged from 0.0 LogMAR to hand movements (mean  $0.7 \pm 0.7$ , median 0.4), 8  
204 had no or mild visual impairment, 3 had moderate visual impairment, 1 had severe, and 1 was  
205 blind. Amongst those over 40 years old (11 individuals), BCVA was between 0.2 LogMAR and  
206 LP (mean  $1.4 \pm 1.1$ , median 0.8), 6 had no or mild impairment, and 5 were blind. Asymmetric  
207 BCVA was seen in 10 patients (21%; 4 in the 15-24 years old group, 5 in the 25-40 years old  
208 group, and 1 in the 41 and older subgroup). Analyzed cross-sectionally, there was a significant  
209 association between age and BCVA for both OD ( $p = 0.0002$ ) and OS ( $p < 0.0001$ , Figure 1C).

210 Refractive data was available for 23 patients (49%), with a mean spherical equivalent of  $-1.9 \pm$   
211  $2.8$  diopters (D). Of these, seventeen patients (74%) had a mild myopic and hyperopic spherical  
212 equivalent ( $-3$  to  $+3$  D), and the remaining 6 had moderate and high myopia.

213

#### 214 Clinical examination, color fundus photography and fundus autofluorescence imaging

215 Corneal examination was unremarkable in all cases. Eleven individuals (23%) developed lens  
216 opacities, at a mean age of  $46.3 \pm 15.2$  (range 26 – 73). There was no evidence of high  
217 intraocular pressure or glaucoma in any patient.

218 Ultrawide-field (UWF) pseudocolor fundus and autofluorescence (AF) imaging was available in  
219 39 participants (82%). The most common retinal feature was the presence of well-demarcated  
220 areas of atrophy at the macula, present in 27 patients (57%), from as early as 16 years of age.  
221 Fine hyperautofluorescent dots at the macula and perimacula (not visible on examination or  
222 colour), were also seen in 23 patients (49%, 6 to 60 years of age), and peripheral punched-out  
223 areas of chorioretinal atrophy were present in 21 patients (45%, Table 1) from 23 years of age,  
224 resembling a more extensive choroideremia-like pattern in patients with advanced disease.

225 Eighteen patients did not have pigment deposits (38%, ages 6 to 64), 15 had minimal bone-  
226 spicule-like (BSL) or nummular pigment (32%, 23 to 72 years old), 3 had moderate BSL  
227 deposits (6%, 32 to 42 years old), and 6 had dense, diffuse BSL pigment (13%, 34 to 79 years of  
228 age). By assessing their imaging, 11 patients (23%) had predominantly macular involvement, 11  
229 (23%) had a prevailing peripheral involvement, and 17 (36%) had widespread retinal dystrophy  
230 affecting the posterior pole and the peripheral retina similarly (Figure 2 and Supplementary  
231 Table 2).

232 Common IRD signs were also present in this cohort, such as peripapillary atrophy (seen in 35  
233 patients, 74%) and a hyperautofluorescent ring at the posterior pole, appearing either entirely  
234 inside the temporal vascular arcades (11 patients) or also encompassing the optic disc (5  
235 individuals, Figure 3A). Other less frequent features were hypoautofluorescent nummular dots  
236 that followed the vascular arcades (5 patients, Figure 3B), a mottled retinal AF pattern (3 young  
237 patients, Figure 3C), foveal-sparing maculopathy (3 patients), and abnormal AF patterns, with  
238 preserved mid-peripheral retina (4 patients, Figure 3D).

239

#### 240 Macular OCT analysis

241 Forty-one individuals had macular OCT scans, Heidelberg blue autofluorescence (BAF) and  
242 infrared (IR) imaging (87%).

243 Baseline central macular thickness (CMT) was  $180.2 \pm 36.0 \mu\text{m}$ , at  $30.2 \pm 13.8$  years old (range  
244 7 – 71). ONLT was quantifiable in 33 patients (66 eyes, 80%, 7 – 51 years old), with a mean  
245 value of  $38.7 \pm 24.4 \mu\text{m}$ . Both CMT and ONLT were significantly lower compared to the  
246 unaffected population ( $p < 0.0001$ ). Foveal EZ was present in 28 patients (52 eyes, 7 to 51 years  
247 old) with a mean width of  $350.5 \pm 302.9 \mu\text{m}$ . A subfoveal hyporeflective zone (HRZ) was seen  
248 in 4 eyes (3 patients, 31 to 50 years old). Considering cross-sectional data only, a significant  
249 association was observed between older age and narrower EZ width ( $p = 0.05$ ), and no  
250 significant association was found between age, CMT and ONLT ( $p = 0.08$  and  $0.35$ ). No  
251 significant association was found between BCVA and EZ, CMT and ONLT ( $p = 0.22$  to  $0.65$ ).

252 Subretinal hyperreflective material was the most common finding, present in 36 individuals  
253 (88%, Figure 4), and this was often correlated with fine hyperautofluorescent dots detected with  
254 BAF. These dots were noticed in the posterior pole or surrounding it in 29 patients (71%) and  
255 were not discernible with IR imaging. Other findings were hyperreflective material in the inner  
256 retina (33 individuals, 80%), epiretinal membrane (ERM, in 27 patients, 66%), and cystoid  
257 macular oedema, present in 3 patients (2 with RP and 1 with CORD).

258

### 259 Electrophysiological assessment

260 Thirty patients had electrophysiological assessment (64%, Tables 1 and 2). Sixteen of them  
261 showed a similar degree of DA and LA ERG reduction, in keeping with a moderate to severe rod  
262 and cone photoreceptor dystrophy, including seven (ages 15-30 years) with undetectable DA and  
263 LA ERGs. The ERG findings in others revealed a rod-cone (N=8; age range 14-51 years; median  
264 24 years) or cone-rod (N=3; age range 15-28 years) pattern of dysfunction, with a further three  
265 showing normal or near-normal ERG amplitudes (cases 28-30; Figure 5; ages 26, 45 and 47  
266 years). The LA 30Hz ERG peak times were normal or borderline (equal to 95th percentile) in the  
267 3 cases with preserved ERG amplitudes (MD group). LA 30Hz ERG peak times were worse in  
268 the 13 patients with the smallest detectable ERGs (median delay 13ms) and were normal or  
269 borderline in the remaining ten, including the five oldest patients (ages 45-51 years).



270 Comparison of ISCEV-standard DA 0.01, DA 3 and DA 10 ERG a- and b-waves, LA 30Hz ERG  
271 and LA 3 (single flash) ERG b-waves revealed a high degree of interocular symmetry in  
272 amplitudes (slope = 1.03;  $r^2=0.98$ ) and peak times (slope = 1.02;  $r^2=0.99$ ). Figure 5 summarizes  
273 the electrophysiological findings (right eyes) and patient ages at the time of testing and Figure 6  
274 shows representative recordings.

275 There was no significant correlation between age and the amplitudes of the DA 0.01 ERG, DA  
276 10 ERG a- and b-waves, LA 30Hz ERG or LA 3 ERG b-waves or the peak times of the LA 30Hz  
277 ERG. Pattern ERG P50 was undetectable in 24 patients in keeping with severe macular  
278 dysfunction. Of the remaining 6 cases, including two with MD, the PERG P50 component was  
279 reduced by 45-60% (Figure 5A). Flash ERGs in the five-year old child tested with skin  
280 electrodes showed only residual photopic ERGs, markedly subnormal scotopic bright flash ERGs  
281 and an undetectable PERG P50 component (data not shown), consistent with a photoreceptor  
282 dystrophy with severe macular involvement.

283 Follow-up ERG data were available in three patients (Supplementary Figure 7), showing rod-  
284 cone (ID 21) or cone-rod (IDs 19 and 20) patterns of dysfunction at baseline. In patient 21 there  
285 was marked reduction in all detectable ERG components and a 15ms increase in LA30Hz ERG  
286 peak time over a 13-year period (from age 14-27 years). In patient 19 there was rapid reduction  
287 in DA ERG amplitudes with no significant change in LA ERG amplitudes or peak times (from  
288 age 15-16 years). In patient 20 there was significant reduction in all ERG components with  
289 borderline (2ms) worsening in LA 30Hz ERG peak time (from age 18 to 23 years).

290

### 291 Longitudinal analysis

292 Mean follow up time was  $9.1 \pm 7.4$  years (range 0-21). Follow up BCVA was available in 34  
293 (72%) patients and was  $1.5 \pm 1$  LogMAR OD (median 1.65) and  $1.65 \pm 1.05$  OS (median 2.28,  
294 mean age  $42 \pm 14$  years). The rate of BCVA decline was 0.08 LogMAR (4 letters)/year and there  
295 was a significant difference between baseline and follow up BCVA ( $p < 0.0001$ ).

296 During the first five-years of follow up ( $n=34$ ), the rate of BCVA decline was 0.08 LogMAR (4  
297 letters)/year for both the overall cohort and for the group aged 15-24 years old, 0.13 (6.5  
298 letters)/year for the 25-40 years old group, and 0.04 (2 letters)/year for the 41 and older



299 subgroup. During years 5-10 of follow up (n=23), the rate was 0.09 LogMAR (4.5 letters)/year  
300 for the 25-40 years old group, and 0.05 (2.5 letters)/year for the older subgroup. Lastly, during  
301 the period of 10-15 years of follow up (n=12), the group aged 25-40 years old had a decrease rate  
302 of 0.08 LogMAR (4 letters)/year, and the older subgroup of 0.06 (3 letters)/year.

303 Twenty-five patients out of the 34 with longitudinal data (74%) had a decrease in BCVA of 15  
304 ETDRS letters or more over follow up in at least one eye, 20 (59%) during the first 5 years since  
305 their baseline visit. Seventeen patients (50%) progressed to more advanced WHO categories of  
306 visual impairment over follow up, 11 (32%) of whom became blind.

307 Thirty-three individuals had longitudinal macular OCT scans during a period of up to 12 years  
308 (mean of  $6.4 \pm 3.5$ ). Longitudinal analysis demonstrated significant differences between baseline  
309 and follow up CMT ( $p < 0.0001$ ,  $-5.9 \mu\text{m}/\text{year}$ ), ONLT ( $p < 0.0001$ ,  $-2.7 \mu\text{m}/\text{year}$ ), and EZ width  
310 ( $p < 0.0001$ ,  $-34.5 \mu\text{m}/\text{year}$ ). Two thirds of the patients had decreased CMT over follow up and  
311 one third had increased values. The ONL was noticed to be isointense to contiguous layers, less  
312 distinct and unable to be quantified in 9 patients over the follow-up period. The association  
313 between BCVA and EZ, CMT and ONLT remained not significant ( $p = 0.16$  to  $0.57$ ). Fifteen out  
314 of 16 patients with a hyperautofluorescent ring at the posterior pole had a follow up assessment.  
315 In 9 patients the ring grew in diameter, in 5 it became smaller, and in one it faded.

316

### 317 Molecular Genetics

318 All patients had biallelic/multiple rare variants in *CERKL*, and their molecular characteristics are  
319 listed in Supplemental Table 3. Thirty-two patients had homozygous variants. Twenty-three  
320 different variants were present in our cohort; 7 nonsense, 6 splice-site alteration (3 canonical  
321 splice site, 2 intronic deletions, and 1 intronic change), 6 missense, 2 small deletions/frameshift,  
322 and 2 large deletions. Eight were classified as pathogenic, 8 as likely pathogenic, and 8 as VUS.  
323 Nine (39%) were novel variants, and 14 were previously reported elsewhere. Schematic  
324 representation of these detected variants and the status of evolutionary conservation are  
325 presented (Figure 8 and Supplementary figure 9).

326 The most common variant was p.(Arg283Ter), encountered in 17 homozygous patients and in 6  
327 presumed compound heterozygotes. The second most frequent variant was p.(Arg106Ser),

328 present in 4 homozygotes and 2 compound heterozygotes. Thirty-six participants (77%) had DN  
329 genotypes (nonsense, frameshift, splicing, or exon deletion), 4 were compound heterozygous of a  
330 null and a missense variant, and 7 had two missense changes (11 NDN, Supplementary Table 3).  
331 One patient (ID 30) was homozygous for a VUS and had a phenotype that matched *CERKL*-  
332 retinopathy, hence he was included in the cohort and considered as a likely *CERKL*-associated  
333 case.

334

### 335 Genotype-phenotype correlation analysis

336 Fifteen out of the 16 patients with childhood-onset symptoms had DN genotypes. The mean age  
337 of onset in individuals with NDN variants was  $22.8 \pm 9.3$  years (median 18.5), and  $19.5 \pm 8.7$  for  
338 DN (median 19,  $p = 0.36$ ). At baseline, the 5 patients aged 15-24 years with moderate visual  
339 impairment harbored DN variants; the 5 patients aged 25-40 years with moderate visual  
340 impairment or worse were DN, and amongst those over 40 years old, 4 out of 5 blind were DN.  
341 The rate of VA decline was  $0.07 \pm 0.07$  LogMAR/year in the NDN group and  $0.08 \pm 0.08$   
342 LogMAR/year in the DN group. There were no significant differences between DN and NDN  
343 groups regarding rate of progression, and no specific variant was found to be consistently milder  
344 than the others or associated with a particular phenotype.

345 Regarding fundus features, 8 out of 11 (73%) NDN patients had no pigment deposits, and the  
346 remaining 3 had minimal BSL pigment (age 20 to 72). In contrast, among 25% of patients with  
347 DN genotypes ( $n=28$  with UWF imaging), 8 had no pigment deposits (29%, ages 14 to 64), 10  
348 had minimal pigment, 3 had moderate, and 4 had dense/diffuse pigmentary changes (aged 34 to  
349 79 years old). The presence of a ring of increased AF at the posterior pole was noted in 50% of  
350 DN patients (14/28) and 27% of NDN (3/11).

351 Concomitant peripheral and macular circumscribed areas of atrophy were observed in 4 patients  
352 with NDN genotypes, versus in 15 with DN genotypes. The DN subgroup included 6 patients  
353 with more marked central retinal involvement, 13 with similar central and peripheral  
354 involvement, and 9 with the peripheral retina mostly affected. The NDN group consisted of 5  
355 patients who had a predominantly macular involvement, 4 who had similar central and peripheral  
356 involvement, and 2 who had a more peripheral degeneration (Table 2).

357

358 The ERG abnormalities were most severe (DA10 ERG a-wave reduction) in 18 of 21 with a DN  
359 genotype, including 7 with undetectable ERGs. The 9 individuals harboring NDN variants  
360 included 3 of the 5 oldest subjects tested, with 3 of the 5 mildest ERG phenotypes (2 with MD;  
361 Figure 5). Of the 10 patients with normal or borderline LA30Hz ERG peak times, 6 harbored  
362 NDN variants.

363 There was significant concordance of phenotype within families, with seven sets of siblings  
364 displaying similar phenotypes, including degree of pigment deposits and atrophic patches, and a  
365 similar age of onset ( $\pm 5$  years). Interestingly, three young siblings were diagnosed with RP and  
366 had a comparable presentation, whereas their mother had a CORD phenotype (IDs 8, 13, 40 and  
367 41). No correlations were found regarding variants' location or gene domain and phenotype.

368

## 369 **Discussion**

370 This study examines the detailed clinical and functional phenotype in the largest cohort of  
371 genetically characterized patients with *CERKL*-associated retinopathy to date. The description of  
372 the uncommon MD presentation is extended. Nine novel disease-causing variants are identified,  
373 and genotype-phenotype correlations examined. Comprehensive retinal imaging, quantitative  
374 electrophysiological and natural history data are detailed, aimed at facilitating the diagnosis and  
375 establishing rates of disease progression, to inform future patient management.

376 *CERKL*-retinopathy has been categorized to date as either presenting as RCD or CORD, even in  
377 patients with the same genotype, excluding possible genotype-phenotype correlations.<sup>15,17,25</sup> It is  
378 of note, that detailed review of our cohort has demonstrated that overlapping features of both  
379 categories were present concurrently in a large proportion of patients; with early drop in acuity  
380 and central visual field loss in patients with 'RCD', and nyctalopia and peripheral visual field loss  
381 as presenting symptoms in individuals with 'CORD' (Supplementary Table 4).<sup>15,17,25</sup> It appears  
382 that unlike the majority of other genetic causes of IRD, the boundaries between the two  
383 categories are not as clear, with many patients presenting with both cone and rod-related  
384 symptoms simultaneously (36% per imaging, 34% per ERG, and 34 % per initial symptoms).

385 This has been noted in previous reports of smaller cohorts, but is extended and confirmed in our  
386 larger, genetically and ethnically diverse cohort (Supplementary Table 2).<sup>5,17</sup>

387 The rate of VA decline was high compared to other genotypes (*CRBI*-RP 0.07 LogMAR/year  
388 and *RPGR*(ORF15) 0.02/year),<sup>24,26</sup> and even faster in the 25-40 year old group, suggesting  
389 preserved acuity until the third decade of life, followed by a more rapid decline. Yet, VA  
390 encompassed an incredibly broad range both at baseline and during follow-up, with over 50% of  
391 individuals above 40 years old still having no or mild visual impairment as per the WHO  
392 classification; indicating the variable prognosis associated with *CERKL*-retinal dystrophy.  
393 Asymmetric BCVA was seen in 21% of patients at baseline, which is high compared to other  
394 conditions including Stargardt disease (reported at 8%).<sup>27</sup> If a less stringent measure of VA  
395 asymmetry was applied, a yet greater asymmetry would be noted.

396 Macular hyperautofluorescent punctate lesions visible with BAF were found to correspond with  
397 subretinal hyperreflective material on OCT (Figure 4A & B). This feature was also highlighted  
398 by Sengillo *et al.*,<sup>28</sup> who described them as becoming denser as the disease progressed. In our  
399 cohort, we observed two scenarios over time; one in which the hyperautofluorescent lesions were  
400 replaced by hypoautofluorescence and increased atrophy, and another where indeed these dots  
401 increased in number as the atrophy grew (Figure 4C & D). The fact that these lesions are excited  
402 by 488 nm blue light could indicate that they are at least partly composed by  
403 lipofuscin/lipofuscin-like and N-retinylidene-N-retinylethanolamine (A2E)/A2E-related material,  
404 suggesting that they may correspond to photoreceptor debris resultant from *CERKL*-mediated  
405 decreased RPE phagocytosis.<sup>29,30</sup> Similar dots/subretinal debris-material has been reported in  
406 *MERTK*-retinopathy,<sup>31</sup> a dysfunctional phagocytosis-associated dystrophy.

407 A novel, less-common finding in our cohort is the presence of a large hyperautofluorescent ring  
408 extending beyond the posterior pole and encompassing the optic disc (5 patients, Supplementary  
409 Table 2 & Figure 3A). This pattern has also been seen in other IRDs including dominant *NR2E3*-  
410 and *EYS*-related retinopathies, representing in both cases the boundaries between affected and  
411 unaffected retina.<sup>32,33</sup> Yet, in *NR2E3*, this ring grew centrifugally towards the mid-periphery and  
412 in *EYS* it shrunk over time. In our case, the ring grew and became less well-defined in the  
413 majority of the patients, in keeping with *NR2E3*-IRD. Other features to consider in *CERKL*-  
414 retinopathy are peripapillary atrophy which, although not pathognomonic of *CERKL*, may be

415 useful when distinguishing between this gene and *ABCA4* (classically associated with  
416 peripapillary sparing);<sup>34</sup> and ERM with inner retinal wrinkling. The latter was reported in up to  
417 22.8% of patients with RP; however, in our cohort it was seen in nearly three times that  
418 proportion, and in previous series it affected 2 and 3 out of 6 patients.<sup>15,35,36</sup>

419 The electrophysiological profiles in our series were diverse, with 16/30 having a similar degree  
420 of rod and cone photoreceptor dysfunction and most others (8/30) having a rod-cone pattern of  
421 dysfunction. A minority had CORD and MD (3/30 each), in keeping with previous reports of  
422 ERG variability.<sup>5,10,15</sup> There was overlap in ERG phenotypes associated with DN and NDN  
423 genotypes, but a large majority of nullizygous cases had severe generalized photoreceptor  
424 dysfunction, including the youngest individuals. Those with NDN variants included two older  
425 patients with MD (normal ERG) and most of those with relatively mild retinal dysfunction. An  
426 apparent disconnect between ERG and initial symptoms was seen in 4 patients with RCD per  
427 electrophysiology and initial symptoms of opposite phenotypes (CORD and MD; ID 6A, 10C,  
428 23, & 24), all with equal central and peripheral involvement in imaging. Although in some of  
429 these cases the symptoms may evolve to eventually match the ERG phenotype (ERG functioning  
430 as a predictor, as in *ABCA4*-Stargardt disease),<sup>37</sup> in others the ERG may represent an additional  
431 piece of information that characterizes the patients disease, not necessarily matching the  
432 remaining assessments. The lack of clear correlation between the ERG findings, symptoms, and  
433 retinal imaging (Supplementary Table 2), highlights the need for comprehensive phenotypic  
434 assessment.

435 In our study, p.(Arg283Ter) was the most common variant, in concordance with other reports.<sup>17</sup>  
436 The second most common change, p.(Arg106Ser), was described to possibly lead to cellular  
437 death given its highly conserved location within the protein, functioning as a null allele despite  
438 being missense.<sup>38</sup> We have demonstrated a degree of genotype-phenotype correlation, with  
439 patients harboring DN genotypes displaying signs of increased severity when compared to those  
440 with NDN, including younger age of onset, greater visual impairment, more retinal pigmentary  
441 disturbance with often concomitant peripheral and macular circumscribed atrophic lesions, and  
442 more profound ERG dysfunction. However, we also noted overlap between the 2 genotypic  
443 groups and the aforementioned signs of disease severity - in keeping with the previously well-

444 documented heterogeneity of IRDs. Investigation of further *CERKL* cases will help to extend our  
445 observations.

446 Over 20 different *CERKL* transcripts have been found in the human retina, due to extensive  
447 alternative splicing and multiple translational start sites, with preferential isoform expression in  
448 different cells.<sup>39,40</sup> A double knockout zebrafish model found that rods degenerated earlier and  
449 more significantly than cones,<sup>30</sup> yet a knockdown-knockout mouse model found cones to be  
450 more affected.<sup>40</sup> The models agreed on photoreceptor outer segment abnormalities, accumulating  
451 in the interphotoreceptor matrix and becoming very long and disorganized, suggestive of  
452 decreased RPE phagocytosis.<sup>30,40</sup> *CERKL* has recently been characterized as an oxidative stress  
453 protective gene within RPE cells, hence decreased RPE function may occur in its absence.<sup>14</sup> It  
454 is possible that different disease-causing variants have greater impact on certain transcripts  
455 preferentially expressed in cones versus rods, or equally in both, resulting in the different  
456 phenotypes and symptoms observed in our study. Furthermore, certain variants could affect how  
457 *CERKL* interacts with other proteins, interfering with its complex function and regulation,  
458 potentially again with different impacts on rods versus cones. Environmental or other genetic  
459 modifying factors may also play a role in the variable phenotypic presentations. Future  
460 functional studies may shed further light on this topic.

461 Given the link between *CERKL* and oxidative stress, antioxidant therapeutic approaches might  
462 have a positive effect on this condition.<sup>41</sup> These are currently under investigation for other IRD  
463 genes such as *ABCA4*, and if successful might also prove useful for *CERKL* retinopathy.<sup>42,43</sup>  
464 Similarly, slowing down the visual cycle could decrease the generation of A2E and limit the  
465 formation of macular hyperautofluorescent punctate lesions. RBP4-inhibitors could be helpful if  
466 proven beneficial in *ABCA4* Stargardt disease (NCT05244304). *CERKL* would also be a good  
467 candidate for gene therapy, given its reasonable window of opportunity. However, due to its size  
468 not fitting in a regular AAV-vector, it would require alternative vectors or dual vector  
469 technology.

470 Our study limitations include its retrospective nature, the different proportion of individuals in  
471 the DN and NDN subgroups, not all data being available for every individual, and data being  
472 acquired with various methods and protocols. The lack of visual field data reflects the real-world  
473 environment in which clinical data were collected. These are to a large extent offset by the large

474 number of genetically-confirmed patients, their wide age-range and ethnic background, and the  
475 international nature of the cohort.

476 This study represents the largest series of patients with *CERKL*-associated retinopathy to date.  
477 We characterised detailed clinical features, disease progression, estimating the rate of visual  
478 acuity change and providing insights into genotype-phenotype correlations. *CERKL*-retinopathy  
479 has a broad phenotypic spectrum ranging from isolated macular dystrophy to severe generalised  
480 retinal involvement. retinal-wide disease and it often does not follow the classical RCD/CORD  
481 dichotomy. and it can present with both cone- and rod-related symptoms at the outset, associated  
482 with structural and functional signs of both cone and rod dysfunction/loss. In many patients, a  
483 greater rate of central vision loss may occur in the third decade. Common retinal changes  
484 observed in *CERKL*-retinopathy include peripheral and macular circular areas of chorioretinal  
485 atrophy, little to no pigment dispersion, and fine hyperautofluorescent macular dots. The nature  
486 of retinal dysfunction often cannot be inferred reliably from clinical signs or imaging alone,  
487 highlighting the importance of comprehensive phenotyping.

488



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620

621 Figure legends

622 **Figure 1:** Visual acuity in *CERKL*-associated retinal dystrophy. A) Mean visual acuity +  
623 standard deviation (SD) in patients from our cohort at their baseline visit, divided into 3 age  
624 groups. B) WHO visual impairment categories in the same age groups at their baseline visit. C)  
625 Linear regression showing a significant association between age and best corrected visual acuity  
626 at baseline visit, for both OD ( $p = 0.0002$ ) and OS ( $p < 0.0001$ ).

627 **Figure 2:** Fundus features of patients with *CERKL*-associated retinal dystrophy. A) 39-year-old  
628 patient with both macular and peripheral circumscribed patches of chorioretinal atrophy (CRA)  
629 and moderate pigment deposits. The patient had a rod-cone dystrophy on electrophysiological  
630 testing (ERG, ID 10) and reported decreased central vision as her initial symptom, at age 15. B)  
631 21-year-old showing only macular atrophy, a hyperautofluorescent ring at the posterior pole, and  
632 no pigmentary changes. He had a cone-rod dystrophy on ERG (ID 27) and described decreased  
633 central vision and light sensitivity since age 17. C) Retinal images from a 45-year-old patient  
634 with predominantly peripheral involvement, with CRA patches in the periphery and mid-  
635 periphery and diffuse pigment dispersion. She did not have an ERG assessment and mentioned  
636 concurrent decreased central vision and nyctalopia since age 36.

637 **Figure 3.** Less common retinal features found in our *CERKL*-associated retinal dystrophy cohort.  
638 A) Autofluorescence (AF) imaging from a 30-year-old man, with a hyperautofluorescent ring at  
639 the posterior pole, extending beyond the temporal vascular arcades and encompassing the optic  
640 disc. Macular OCT scan showing remnant subfoveal ellipsoid zone (EZ) line. The patient had  
641 equally affected rod and cone ERG responses (ID 22) and was diagnosed with a 'macular  
642 dystrophy' at age 19. B) 32-year-old patient with a hypoautofluorescent pattern that follows the  
643 vascular arcades. Macular OCT scan is positive for outer retinal degeneration and an epiretinal  
644 membrane (ERM). She had equal rod and cone dysfunction on ERG (ID 3) and reported  
645 decreased central vision as her initial symptom, at age 7. C) AF imaging from a 15-year-old  
646 patient uneven autofluorescence and hypoautofluorescent dots inferiorly. Macular OCT shows  
647 minimal subfoveal EZ and outer retinal degeneration. He had equal rod and cone impairment on  
648 ERG (ID 13) and developed decreased central vision at age 13. D) 55-year-old patient with a few  
649 peripheral CRA patches and an abnormal autofluorescence pattern, with a preserved nasal, mid-  
650 peripheral area. Macular OCT shows minimal subfoveal EZ, outer retinal degeneration and

651 ERM. She had a rod-cone dystrophy pattern on ERG (ID 24) and described decreased central  
 652 vision since age 23. E) 9-year-old (ID 41, brother of ID 13 -displayed on C-). There is vessel  
 653 thinning, mild retinal mottling and uneven autofluorescence. Bilateral macular OCT is positive  
 654 for outer retinal loss, with preserved central EZ and overall architecture. He remains  
 655 asymptomatic (last visit at age 11).

656 **Figure 4.** Macular optical coherence tomography (OCT) and Heidelberg imaging. A) Correlation  
 657 of blue fundus autofluorescence (BAF), infrared (IR) and OCT imaging of patient ID 14 at 20  
 658 years of age, with rod-cone dystrophy on ERG and mixed rod and cone symptoms at onset. A  
 659 hyperautofluorescent dot is seen in BAF next to the vessel (green square), hyporeflective in IR  
 660 and subretinal hyperreflective material evident in that location on OCT. B) Correlation of BAF,  
 661 IR and OCT imaging of patient ID 20 at age 27. Here is another example of a  
 662 hyperautofluorescent dot on BAF, hyporeflective on IR and subretinal hyperreflective material in  
 663 that location. C) Fundus autofluorescence images of a patient clinically diagnosed with *CERKL*-  
 664 associated retinitis pigmentosa (ID 43). To the left is the patient at 26 years of age and to the  
 665 right that same patient at age 37. Scattered atrophy is seen in the posterior pole and mid-  
 666 periphery, with hyperautofluorescent fine dots surrounding the posterior pole at an early stage,  
 667 and few dots remaining in the most recent image, these having been largely replaced by  
 668 hypoautofluorescence. Macular OCT scans show a narrowed EZ line and loss of retinal layering  
 669 over follow up. D) Fundus autofluorescence changes in patient ID 20, with an  
 670 electrophysiological diagnosis of cone-rod dystrophy and mixed rod and cone symptoms at  
 671 onset. To the left is the posterior pole at age 18 and to the right at age 33. In this case, the central  
 672 atrophy enlarged, and the hyperautofluorescent dots increased in number/ became more  
 673 noticeable over time. Longitudinal macular OCT shows more profound outer retinal atrophy and  
 674 poor retinal architecture over time.

675 **Figure 5.** Full field ERG and PERG findings summarized in 30 patients tested according the  
 676 ISCEV standard; a) The amplitudes of the DA 10 ERG a-wave, LA 30 Hz ERG, LA 3 ERG b-  
 677 wave and PERG P50 component are plotted against the primary axis as a percentage of the age-  
 678 matched lower limit of the reference range (horizontal broken line), with values arranged in  
 679 ascending order of DA 10 ERG a-wave amplitude for clarity. The LA 30 Hz peak times are  
 680 plotted against the secondary axis as a difference from the age-matched upper limit of the

681 reference range (horizontal dotted line). b) The age of the patients at the time of testing, arranged  
682 in same order as in a). Patients with NDN variants include those numbered 11, 14, 20, 22, 23, 25,  
683 27, 29 and 30 (highlighted with vertical arrows). All DA and LA ERGs and PERGs were  
684 undetectable in patients 1-5, 6 and 7.

685 **Figure 6.** Representative full-field and pattern ERGs from three patients and one control subject  
686 for comparison; a) patient 2 (aged 30 years); b) patient 22 (29 years); c) patient 30 (47 years); d)  
687 representative control (“normal”) recordings. Numbering of patients corresponds to that used in  
688 Figure 5. Data are shown for the right eyes only, as all recordings showed a high degree of inter-  
689 ocular symmetry. Patient traces are superimposed to demonstrate reproducibility. Broken lines  
690 replace blink artefacts for clarity. The full-field ERGs in patients 2 and 22 are consistent with a  
691 rod and cone photoreceptor dystrophy affecting rods and cones similarly, with PERG evidence of  
692 macular involvement, more severe in patient 2. The ERGs in patient 21 are in keeping with a  
693 rod-cone dystrophy, with a detectable but subnormal PERG, indicating macular involvement.  
694 The undetectable PERG in case 30 is consistent with severe macular dysfunction with no full-  
695 field ERG evidence of generalized retinal dysfunction.

696 **Figure 8.** Graphic representation of the *CERKL* gene and protein, with details on functional  
697 domains. *CERKL* (Transcript; NM\_001030311.2, ENST00000339098.9, Uniprot accessions:  
698 Q49MI3) contains 13 exons that encode a 532 amino acid protein containing a disordered region,  
699 nuclear localization signal 1 motif, nuclear localization signal 2 motif, and a diacylglycerol  
700 kinase catalytic domain. Seven nonsense, 6 missense, and 2 small deletion/frameshift variants  
701 detected in this cohort are demonstrated. Novel variants are highlighted by a \*.

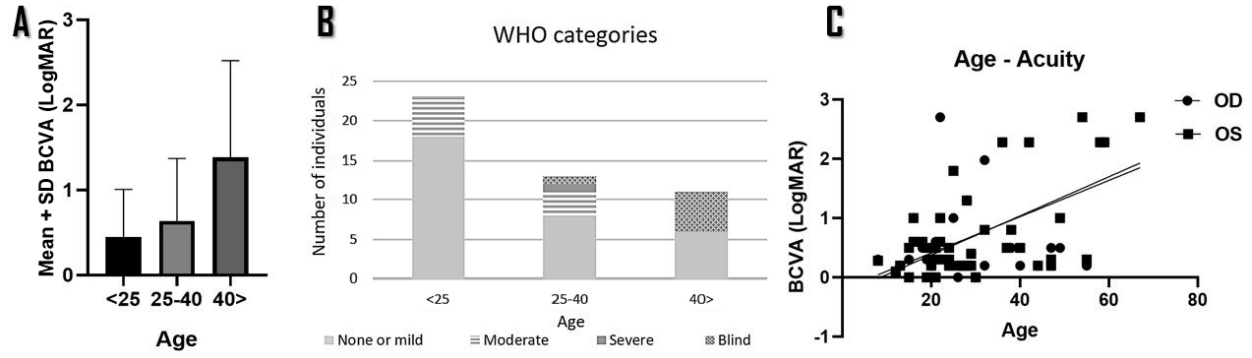
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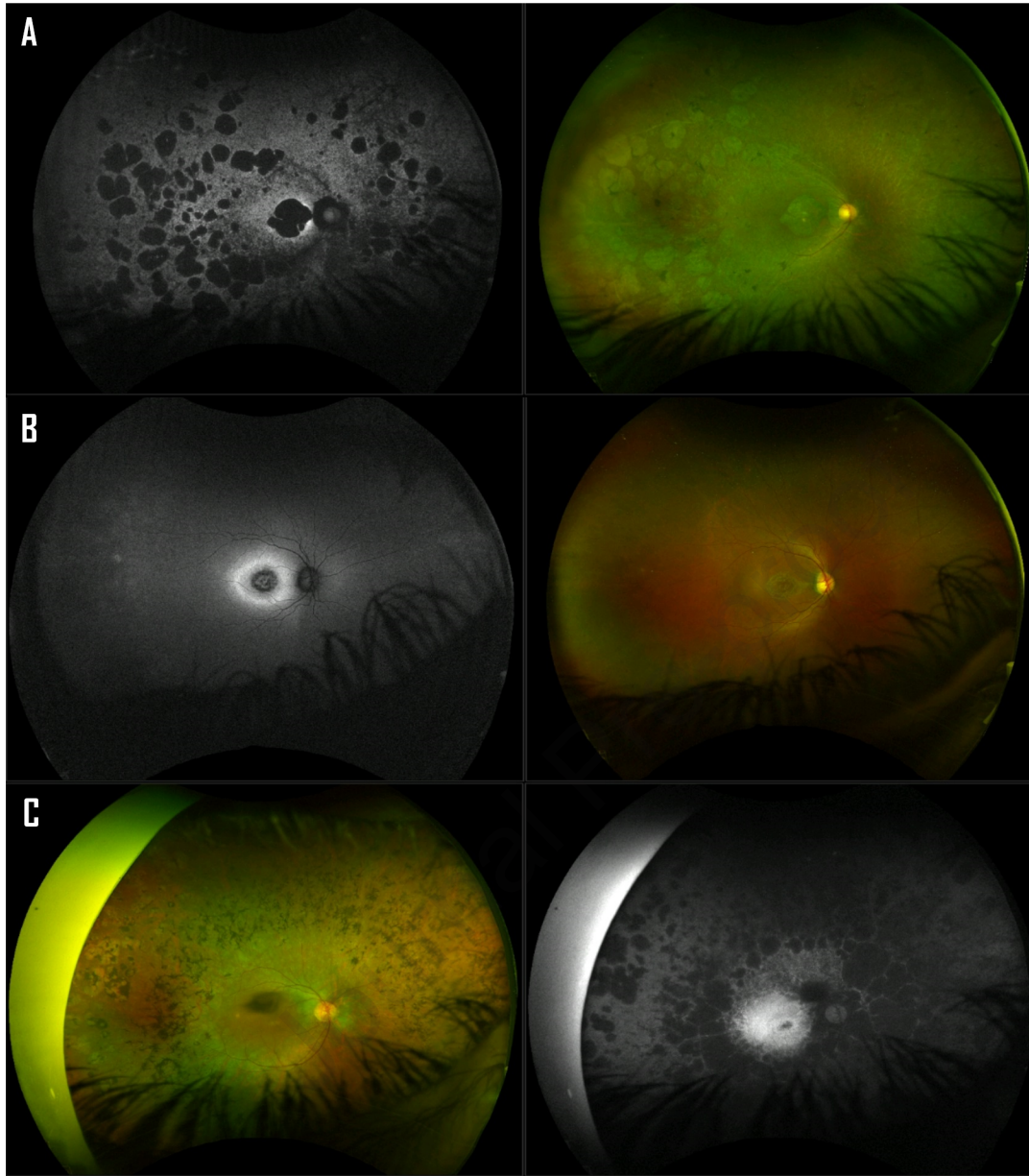
**Table 1.** Clinical Characteristics of Patients with *CERKL*-Associated Retinal Dystrophy

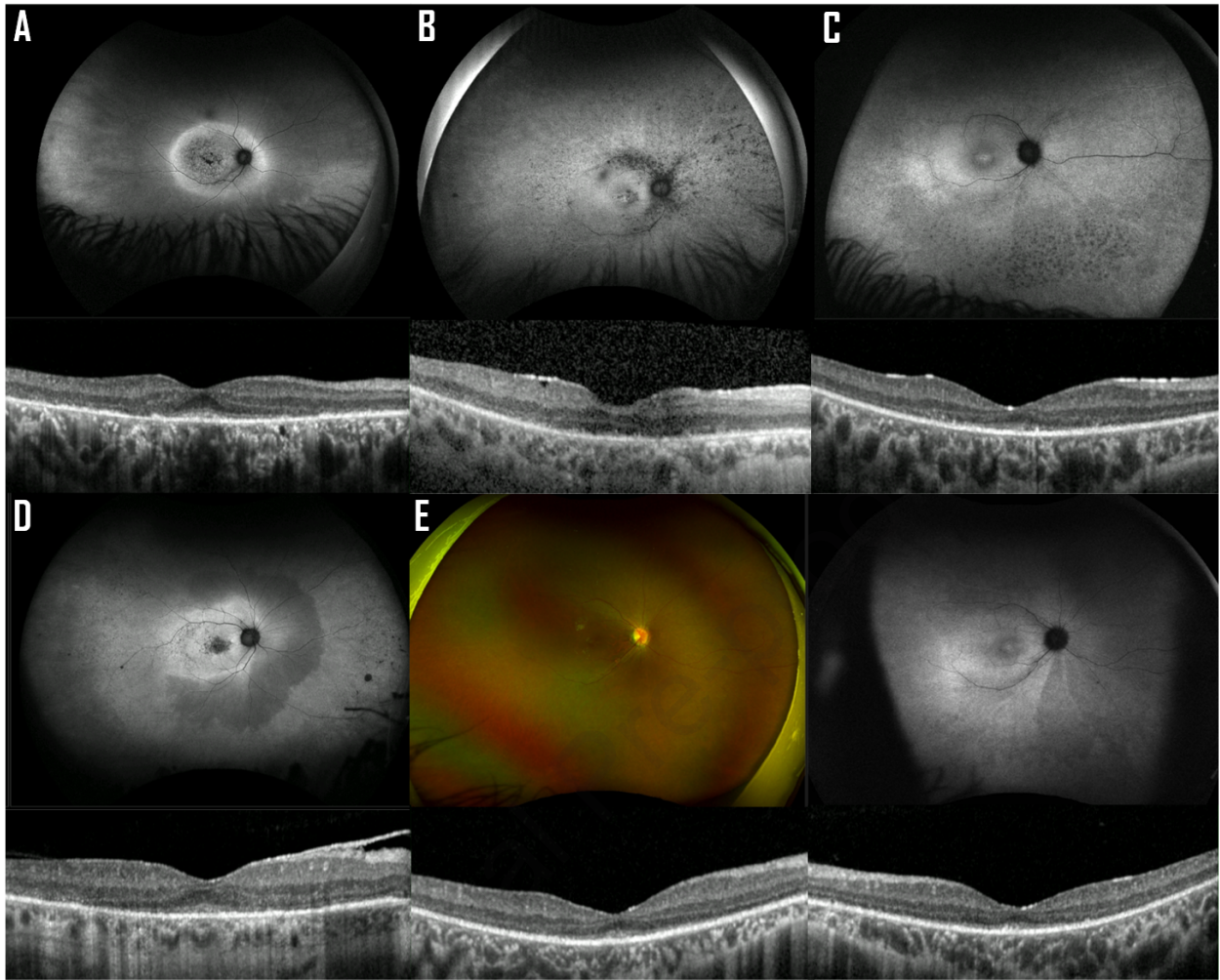
Characteristics	Patients (n=47)
Families	37
Gender (n (%))	
. Female	22 (47)
. Male	25 (53)
Age at first examination (mean $\pm$ SD, years)	29.6 $\pm$ 13.9
Age at last examination (mean $\pm$ SD, years)	39.2 $\pm$ 16.6
Follow-up time (mean $\pm$ SD, years)	9.1 $\pm$ 7.4
Ethnicity (n (%))	34 (72)
. Asian	19 (56)
. White European	13 (38)
. African descent	2 (6)
Age of onset (mean $\pm$ SD, years)	19.8 $\pm$ 9 years
Reported first symptom (n (%))	
. Central vision loss	19 (40)
. Concurrent central vision loss and nyctalopia	16 (34)
. Night blindness	5 (11)
. Others (scotomas, photophobia, colour vision issues)	7 (15)
Spherical equivalent refractive error (mean $\pm$ SD, D)	-1.9 $\pm$ 2.8
Funduscopy examination (n (%))	
. Circumscribed areas of atrophy in the macula	27 (57)
. Fine white dots at the macula and perimacula	23 (49)
. Peripheral punched-out-like areas of chorioretinal atrophy	21 (45)
. Hyperautofluorescent ring at the posterior pole (Robson ring)	11 (23)
. Ring including the optic disc	5 (11)
. Hypoautofluorescent nummular dots that followed the vascular arcades	5 (11)
. Mottled/uneven retinal aspect	3 (6)
. foveal-sparing maculopathy	3 (6)
. Abnormal autofluorescence pattern	4 (9)
Retinal pigment deposits (n (%))	42 (89)
. None	18 (38)
. Minimal	15 (32)
. Moderate	3 (6)
. Dense, diffuse	6 (13)
Most affected retinal area by UWF imaging (n (%))	39 (82)
. Macula	11 (23)
. Periphery	11 (23)
. Similar both	17 (36)
Full-field electroretinography (n (%))	30 (64)
. Similar cone and rod involvement	16 (53)
. Rod-cone pattern	8 (27)
. Cone-rod pattern	3 (10)
. Macular dystrophy (normal full-field ERG)	3 (10)

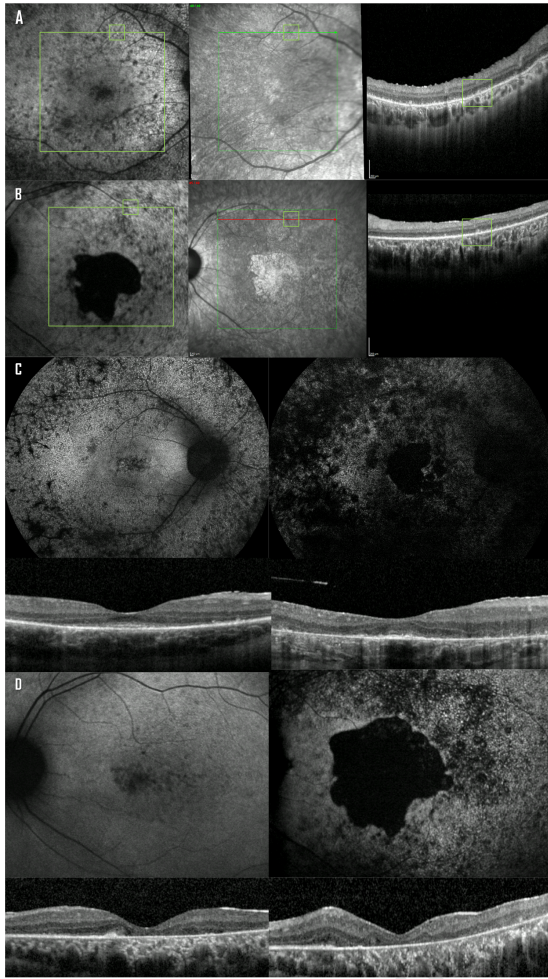
**Abbreviations:** SD: standard deviation; n: number; D: diopters; UWF: ultrawide-field; ERG: electroretinogram



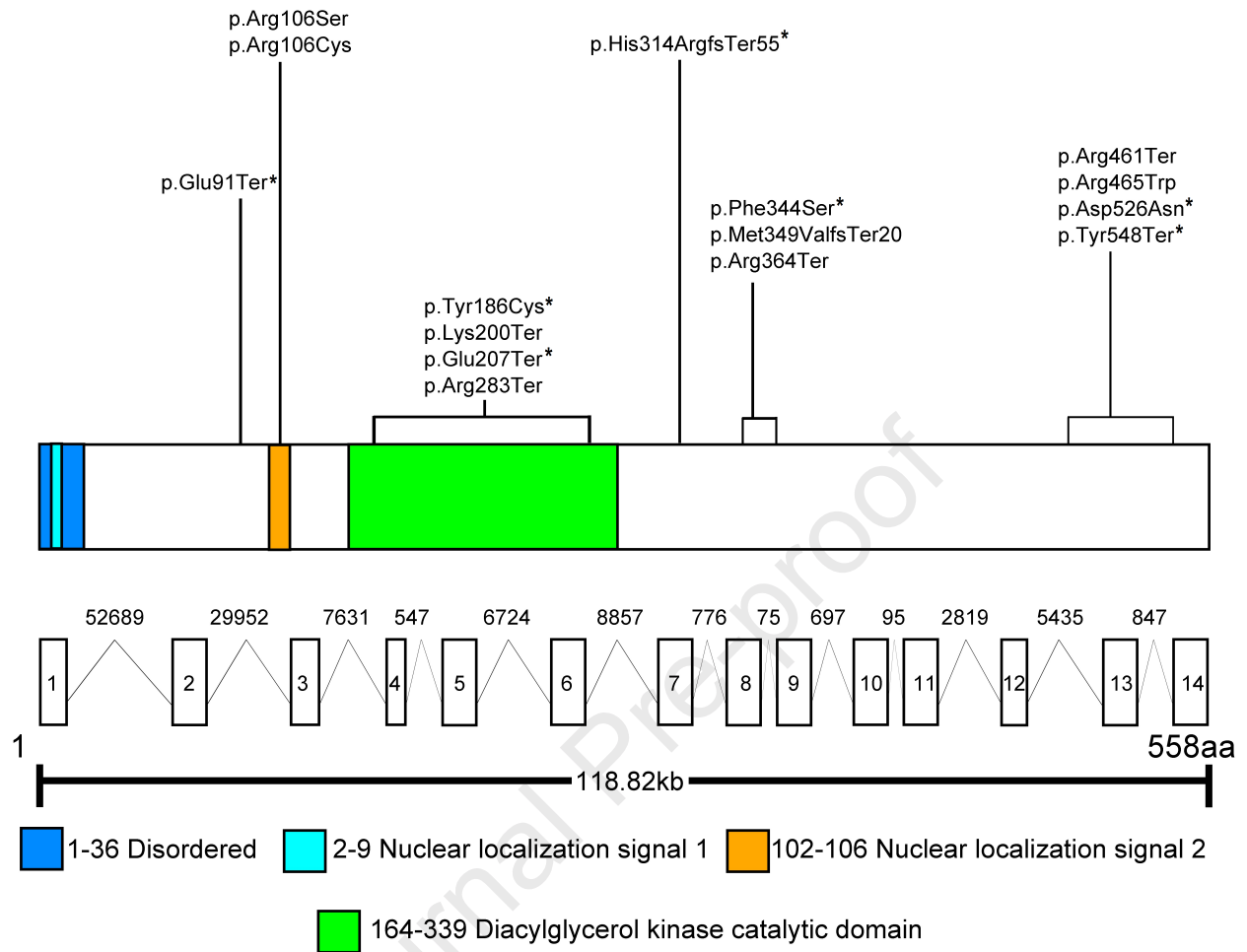




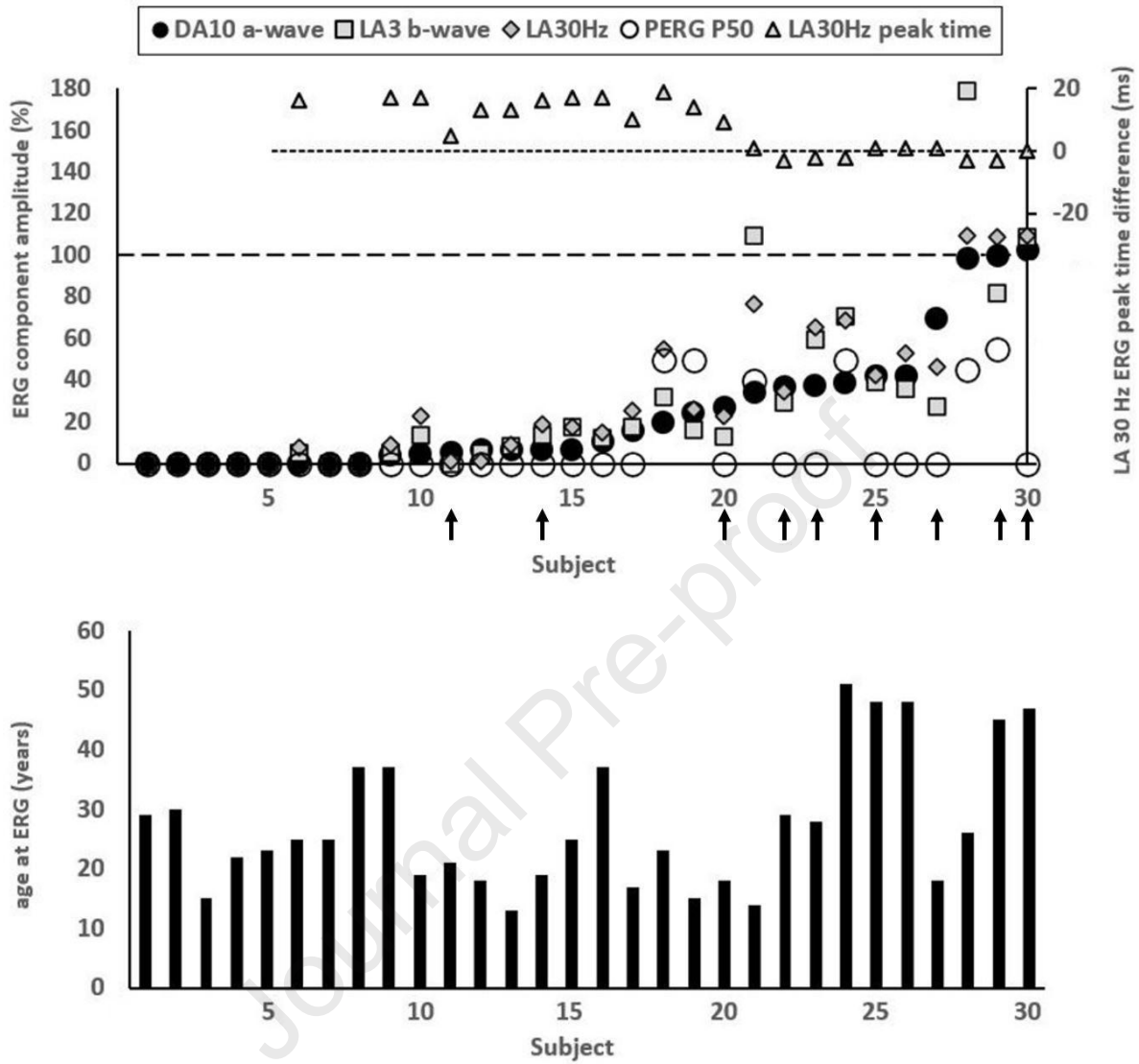


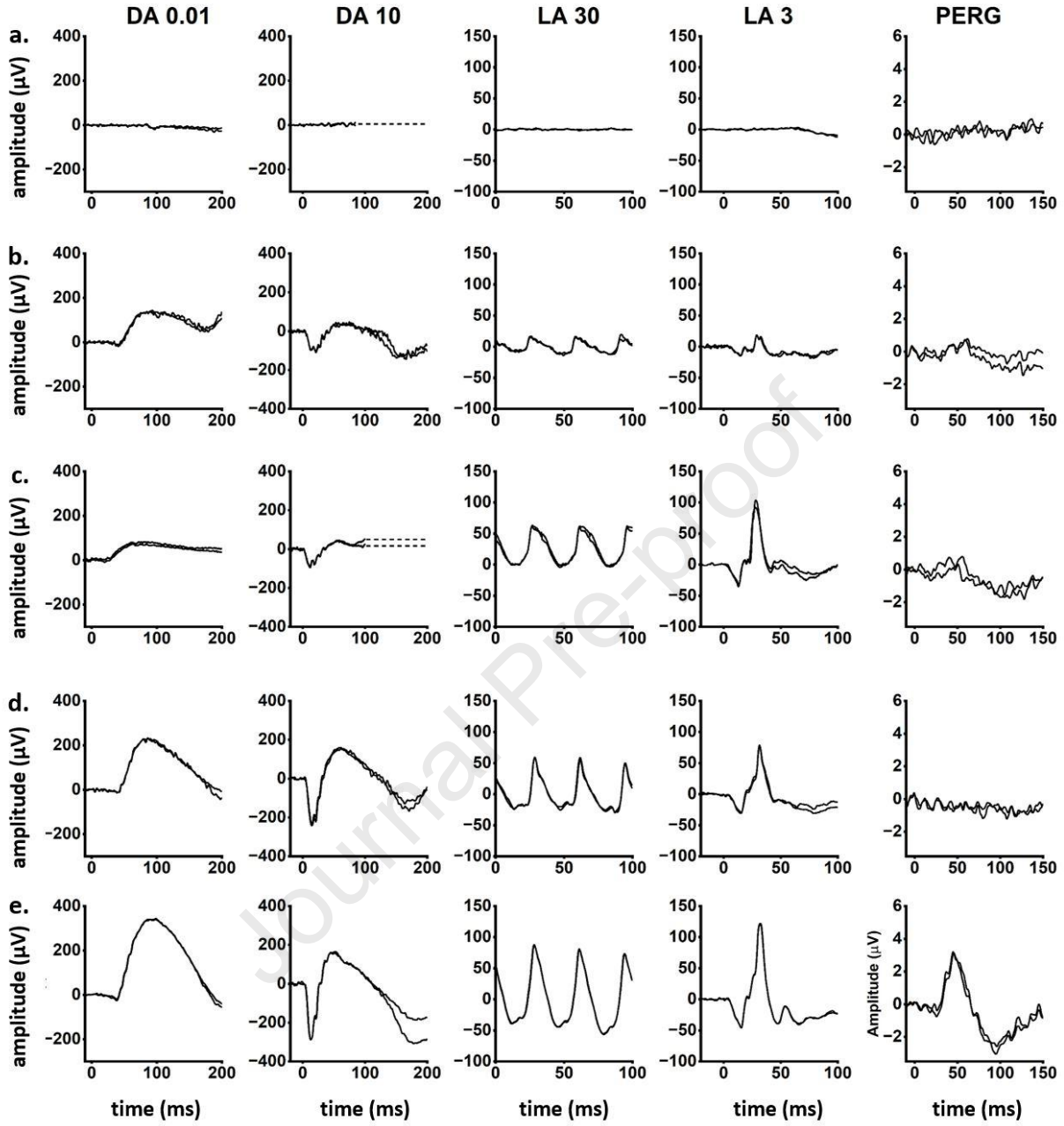


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*CERKL*-retinopathy encompasses isolated macular disease to severe retina-wide involvement, with a range of functional phenotypes, generally not fitting in the rod-cone/cone-rod dichotomy. Nullizygous cases are often more severe.

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